

10/019284

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SEQUENCE LISTING

<110> IZUI, MASAKO
SUGIMOTO, MASAKAZU
KURAHASHI, OSAMU
NAKAMATSU, TSUYOSHI

<120> DNA ENCODING SUCROSE PTS ENZYME II

<130> 217677US0PCT

<140> US 10/019,284

<141> 2002-01-02

<150> JP 11-189512

<151> 1999-07-02

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<170> PatentIn version 3.1

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 Arg Arg Gln His Cys Arg Arg Arg Thr Leu Cys Asn Ala Phe Thr Pro
 20 25 30
 cgt gct caa aga cac caa gga tgt gga tcg cca aag tct gga tga tga 3922
 Arg Ala Gln Arg His Gln Gly Cys Gly Ser Pro Lys Ser Gly
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Arg Arg Ala Arg Arg Cys Gly Ser Cys Phe Gln Arg Thr Arg Arg	
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Asn Leu Gln Arg His Arg Cys Val His Arg Ala Ala Gln Arg Cys Cys	
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ggc taa caa cgc caa ctg gtt cag ccg tgc tgt gaa ggt att ggc gga	4114
Gly Gln Arg Gln Leu Val Gln Pro Cys Cys Glu Gly Ile Gly Gly	
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cat ttt cgt ccc gct gat tcc aat ctt ggt tgg tgg cgg tct gct cat	4162
His Phe Arg Pro Ala Asp Ser Asn Leu Gly Trp Trp Arg Ser Ala His	
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Gly Tyr Gln Gln Cys Val Gly Cys Ala Gly Ser Val Arg Ser Ala Ile	
125 130 135	
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Thr Gly Gly Asp Val Pro Ser Asp Gln Arg Cys Cys Asp Asp Gln	
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Pro Asp Gly Ile Cys Ala Val Arg Val Leu Ala Ser Val Gly Trp Phe	
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His Arg Asn Gln Ala Phe Arg Trp Gln Val Pro Gly Arg Arg His	
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Trp Tyr Gly Asp Gly Val Pro Asn Pro Gly Arg Leu Arg Arg Gly	
185 190 195	
cgc cac cat gac cgc ggg cga aat gcc aat gtg gtc cct gtt tgg ttt	4450
Arg His His Asp Arg Gly Arg Asn Ala Asn Val Val Pro Val Trp Phe	
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Gly Cys Cys Ser Ser Trp Leu Pro Gly His Arg Ala Ser Cys Ala Gly	
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Gly Leu Leu Asp Ser Gly Asn Asp Arg Glu Val Pro Ala Gln Ala Thr	
235 240 245	
cat ggg cac tgc aga ctt cct gat cac ccc agt gtt gac tct gct gct	4594
His Gly His Cys Arg Leu Pro Asp His Pro Ser Val Asp Ser Ala Ala	
250 255 260	
cac cgg ctt cct tac gtt cat tgc tat tgg tcc agc aat gcg ctg ggt	4642

His Arg Leu Pro Tyr Val	His Cys Tyr Trp Ser	Ser Asn Ala Leu Gly	
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ggg tga ctt gct ggc aca	cgg tct gca ggg act	cta tga ttt cgg tgg	4690
Gly Leu Ala Gly Thr	Arg Ser Ala Gly Thr	Leu Phe Arg Trp	
280	285	290	
tcc agt cgg cgg tct gct	ttt cgg tct ggt cta	ctc acc aat cgt tat	4738
Ser Ser Arg Arg Ser Ala	Phe Arg Ser Gly Leu	Leu Thr Asn Arg Tyr	
295	300	305	
cac tgg tct gca cca gtc	ctt ccc gcc aat tga	gct gga gct gtt caa	4786
His Trp Ser Ala Pro Val	Leu Pro Ala Asn	Ala Gly Ala Val Gln	
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cca ggg tgg atc ctt cat	chc gca acc gca tcc	atg gcc aat atc gcg	4834
Pro Gly Trp Ile Leu His	Xaa Ala Thr Ala Ser	Met Ala Asn Ile Ala	
325	330	335	340
cag ggt gca gca tgt ttg	gca gtg ttc ttc cta	gcg aag agt gaa aag	4882
Gln Gly Ala Ala Cys Leu	Ala Val Phe Phe Leu	Ala Lys Ser Glu Lys	
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ctc aag ggc ctt gca ggt	gct tca ggt gtc tcc	gct gtt ctt ggt att	4930
Leu Lys Gly Leu Ala Gly	Ala Ser Gly Val Ser	Ala Val Leu Gly Ile	
360	365	370	
aca gag cct gcg atc ttc	ggt gtg aac ctt cgc	ctg cgc tgg ccg ttc	4978
Thr Glu Pro Ala Ile Phe	Gly Val Asn Leu Arg	Leu Arg Trp Pro Phe	
375	380	385	
tac att ggt atc ggt acc	gca gct atc ggt ggc	gct ttg att gca ctc	5026
Tyr Ile Gly Ile Gly Thr	Ala Ala Ile Gly Gly	Ala Leu Ile Ala Leu	
390	395	400	
ttt gat atc aag gca gtt	gcg ttg ggc gct gca	ggg ttc ttg ggt gtt	5074
Phe Asp Ile Lys Ala Val	Ala Leu Gly Ala Ala	Gly Phe Leu Gly Val	
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ggt tct att gat gct cca	gat atg gtc atg ttc	ttg gtt tgc gcg gta	5122
Val Ser Ile Asp Ala Pro	Asp Met Val Met Phe	Leu Val Cys Ala Val	
425	430	435	
ggt acc ttt gtc atc gca	ttc ggc gca gcg att	gct tat ggc ctt tac	5170
Val Thr Phe Val Ile Ala	Phe Gly Ala Ala Ile	Ala Tyr Gly Leu Tyr	
440	445	450	
ttg gtt cgc cgc aac ggc	agc att gat cca gat	gca acc gct gct cca	5218
Leu Val Arg Arg Asn Gly	Ser Ile Asp Pro Asp	Ala Thr Ala Ala Pro	
455	460	465	
gtg cct gca gga acg acc	aaa gcc gaa gca gaa	gca ccc gca gaa ttt	5266
Val Pro Ala Gly Thr Thr	Lys Ala Glu Ala Glu	Ala Pro Ala Glu Phe	
470	475	480	
tca aac gat tcc acc atc	atc cag gca cct ttg	acc ggt gaa gct atc	5314
Ser Asn Asp Ser Thr Ile	Ile Ile Gln Ala Pro	Leu Thr Gly Glu Ala Ile	

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Ala Leu Ser Ser Val Ser Asp Ala Met Phe Ala Ser Gly Lys Leu Gly	505	510	515	
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Ser Gly Val Ala Ile Val Pro Thr Lys Gly Gln Leu Val Ser Pro Val	520	525	530	
agc gga aag atc gtg gtg gcc ttc cca tct ggt cac gct ttc gca gtc				5458
Ser Gly Lys Ile Val Val Ala Phe Pro Ser Gly His Ala Phe Ala Val	535	540	545	
cgc act aag gct gag gat ggt tcc aat gtg gat atc ttg atg cac att				5506
Arg Thr Lys Ala Glu Asp Gly Ser Asn Val Asp Ile Leu Met His Ile	550	555	560	
ggt ttc gac acc gta aac ctc aac ggc acg cac ttt aac ccg ctg aag				5554
Gly Phe Asp Thr Val Asn Leu Asn Gly Thr His Phe Asn Pro Leu Lys	565	570	575	580
aag cag ggc gat gaa gtc aaa gca ggg gag ctg ctg tgt gaa ttc gat				5602
Lys Gln Gly Asp Glu Val Lys Ala Gly Glu Leu Leu Cys Glu Phe Asp	585	590	595	
att gat gcc att aag gct gca ggt tat gag gta acc acg ccg att gtt				5650
Ile Asp Ala Ile Lys Ala Ala Gly Tyr Glu Val Thr Thr Pro Ile Val	600	605	610	
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Val Ser Asn Tyr Lys Lys Thr Gly Pro Val Asn Thr Tyr Gly Leu Gly	615	620	625	
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Glu Ile Glu Ala Gly Ala Asn Leu Leu Asn Val Ala Lys Lys Glu Ala	630	635	640	
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Val Pro Ala Thr Pro	645			
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20 25 30

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 35 40 45

Pro Ala Gln Ala Thr His Gly His Cys Arg Leu Pro Asp His Pro Ser
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Ser Asn Ala Leu Gly Gly
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Ala	Lys	Ser	Glu
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Lys	Leu	Lys	Gly
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Leu	Ala	Gly	Ala
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Ser	Gly	Val	Ser
Ala	Val	Leu	Gly
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Ile	Thr	Glu	Pro
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Ala	Ile	Phe	Gly
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Val	Asn	Leu	Arg
Leu	Arg	Trp	Pro
65			
Phe	Tyr	Ile	Gly
70			
Ile	Gly	Ile	Gly
75			
Thr	Ala	Ala	Ile
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Gly	Gly		
Ala	Leu	Ile	Ala
85			
Leu	Phe	Asp	Ile
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Lys	Ala	Val	Ala
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Leu	Gly	Ala	Ala
Gly	Phe	Leu	Gly
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Val	Val	Ser	Ile
105			
Asp	Ala	Pro	Asp
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Leu	Val	Cys	Ala
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Val	Val	Thr	Phe
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Ala	Tyr	Gly	Leu
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Tyr	Leu	Val	Arg
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Arg	Arg	Asn	Gly
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Ser	Ile	Asp	Pro
Ala	Thr	Ala	Ala
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Gly	Thr	Thr	Lys
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Ala	Glu	Ala	Glu
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Ser	Gly	Lys	Leu
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Gly	Ser	Gly	Val
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Ala	Ile	Val	Pro
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Thr	Lys	Gly	Gln
Leu	Val	Ser	Pro
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Val	Ser	Gly	Lys
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Ile	Val	Val	Ala
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Phe	Pro	Ser	Gly
His	Ala	Phe	Ala
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Val	Arg	Thr	Lys
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Ala	Glu	Asp	Gly
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Ser	Asn	Val	Asp
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Leu Cys Glu Phe Asp Ile Asp Ala Ile Lys Ala Ala Gly Tyr Glu Val
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Thr Thr Pro Ile Val Val Ser Asn Tyr Lys Lys Thr Gly Pro Val Asn
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